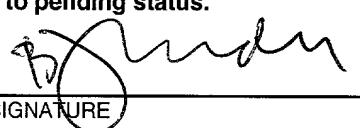


FORM PTO-1390 (REV 11-98)	U.S. DEPARTMENT OF COMMERCE PATENT AND TRADEMARK OFFICE	ATTORNEY'S DOCKET NUMBER 117-347
TRANSMITTAL LETTER TO THE UNITED STATES DESIGNATED/ELECTED OFFICE (DO/EO/US) CONCERNING A FILING UNDER 35 U.S.C. 371		U.S. APPLICATION NO. (If known, see 37 C.F.R. 1.5) 09/787633 <small>Unknown</small>
INTERNATIONAL APPLICATION NO. PCT/GB99/03180	INTERNATIONAL FILING DATE 22 September 1999	PRIORITY DATE CLAIMED 22 September 1998
TITLE OF INVENTION TREATMENT OF INFECTION		
APPLICANT(S) FOR DO/EO/US WILSON et al.		
Applicant herewith submits to the United States Designated/Elected Office (DO/EO/US) the following items and other information:		
1. <input checked="" type="checkbox"/> This is a FIRST submission of items concerning a filing under 35 U.S.C. 371. 2. <input type="checkbox"/> This is a SECOND or SUBSEQUENT submission of items concerning a filing under 35 U.S.C. 371. 3. <input checked="" type="checkbox"/> This is an express request to begin national examination procedures (35 U.S.C. 371(f) at any time rather than delay examination until the expiration of the applicable time limit set in 35 U.S.C. 371(b) and PCT Articles 22 and 39(1). 4. <input checked="" type="checkbox"/> A proper Demand for International Preliminary Examination was made by the 19 th month from the earliest claimed priority date. 5. A copy of the International Application as filed (35 U.S.C. 371(c)(2)). a. <input type="checkbox"/> is transmitted herewith (required only if not transmitted by the International Bureau). b. <input checked="" type="checkbox"/> has been transmitted by the International Bureau. c. <input type="checkbox"/> is not required, as the application was filed in the United States Receiving Office (RO/US). 6. <input type="checkbox"/> A translation of the International Application into English (35 U.S.C. 371(c)(2)). 7. <input checked="" type="checkbox"/> Amendments to the claims of the International Application under <u>PCT Article 34</u> . a. <input type="checkbox"/> are transmitted herewith (required only if not transmitted by the International Bureau). b. <input checked="" type="checkbox"/> have been transmitted by the International Bureau. c. <input type="checkbox"/> have not been made; however, the time limit for making such amendments has NOT expired. d. <input type="checkbox"/> have not been made and will not be made. 8. <input type="checkbox"/> A translation of the amendments to the claims under PCT Article 19 (U.S.C. 371(c)(3)). 9. <input type="checkbox"/> An oath or declaration of the inventor(s) (35 U.S.C. 371(c)(4)). 10. <input type="checkbox"/> A translation of the annexes to the International Preliminary Examination Report under PCT Article 36 (35 U.S.C. 371(c)(5)). Items 11. To 16. Below concern document(s) or information included: 11. <input checked="" type="checkbox"/> An Information Disclosure Statement under 37 C.F.R. 1.97 and 1.98. 12. <input type="checkbox"/> An assignment document for recording. A separate cover sheet in compliance with 37 C.F.R. 3.28 and 3.31 is included. 13. <input checked="" type="checkbox"/> A FIRST preliminary amendment. <input type="checkbox"/> A SECOND or SUBSEQUENT preliminary amendment. 14. <input type="checkbox"/> A substitute specification. 15. <input type="checkbox"/> A change of power of attorney and/or address letter. 16. <input checked="" type="checkbox"/> Other items or information attached: IDS; PTO 1449 with copies of listed documents; paper & computer readable copies of Sequence Listing; Letter re: Sequence Listing <input checked="" type="checkbox"/> This application is entitled to "Small entity" status. <input type="checkbox"/> "Small entity" statement attached.		

U.S. APPLICATION NO. (If known, see 37 C.F.R. 1.5) Unknown 09/787633		INTERNATIONAL APPLICATION NO. PCT/GB99/03180		ATTORNEY'S DOCKET NUMBER 117-347	
17. <input checked="" type="checkbox"/> The following fees are submitted:				CALCULATIONS PTO USE ONLY	
BASIC NATIONAL FEE (37 C.F.R. 1.492(a)(1)-(5): -- Neither international preliminary examination fee (37 C.F.R. 1.482) nor international search fee (37 C.F.R. 1.445(a)(2)) paid to USPTO and International Search Report not prepared by the EPO or JPO\$1000.00 -- International preliminary examination fee (37 C.F.R. 1.482) not paid to USPTO but International Search Report prepared by the EPO or JPO.....\$860.00 -- International preliminary examination fee (37 C.F.R. 1.482) not paid to USPTO but international search fee (37 C.F.R. 1.445(a)(2)) paid to USPTO\$710.00 -- International preliminary examination fee paid to USPTO (37 C.F.R. 1.482) but all claims did not satisfy provisions of PCT Article 33(1)-(4).....\$690.00 -- International preliminary examination fee paid to USPTO (37 C.F.R. 1.482) and all claims satisfied provisions of PCT Article 33(1)-(4).....\$100.00 <div style="text-align: right;">ENTER APPROPRIATE BASIC FEE AMOUNT =</div>				<div style="border: 1px solid black; padding: 2px;">\$ 860.00</div>	
Surcharge of \$130.00 for furnishing the oath or declaration later than <input type="checkbox"/> 20 <input checked="" type="checkbox"/> 30 months from the earliest claimed priority date (37 C.F.R. 1.492(e)).				<div style="border: 1px solid black; padding: 2px;">\$ 130.00</div>	
CLAIMS	NUMBER FILED	NUMBER EXTRA	RATE		
Total Claims	8	-20 =	0	X	\$18.00
Independent Claims	4	-3 =	1	X	\$80.00
MULTIPLE DEPENDENT CLAIMS(S) (if applicable)					\$270.00
TOTAL OF ABOVE CALCULATIONS =					\$ 1070.00
Reduction by 1/2 for filing by small entity, if applicable. Small entity status must also be asserted. (Note 37 C.F.R. 1.9, 1.27, 1.28).					535.00
SUBTOTAL =					\$ 535.00
Processing fee of \$130.00, for furnishing the English Translation later than <input type="checkbox"/> 20 <input type="checkbox"/> 30 months from the earliest claimed priority date (37 C.F.R. 1.492(f)).					0.00
TOTAL NATIONAL FEE =					\$ 535.00
Fee for recording the enclosed assignment (37 C.F.R. 1.21(h)). The assignment must be accompanied by an appropriate cover sheet (37 C.F.R. 3.28, 3.31). \$40.00 per property					0.00
Fee for Petition to Revive Unintentionally Abandoned Application (\$1240.00 - Small Entity = \$620.00)					0.00
TOTAL FEES ENCLOSED =					\$ 535.00
				Amount to be:	
				refunded	\$
				Charged	\$
a. <input checked="" type="checkbox"/> A check in the amount of \$535.00 to cover the above fees is enclosed. b. <input type="checkbox"/> Please charge my Deposit Account No. 14-1140 in the amount of \$_____ to cover the above fees. A duplicate copy of this form is enclosed. c. <input checked="" type="checkbox"/> The Commissioner is hereby authorized to charge any additional fees which may be required, or credit any overpayment to Deposit Account No. 14-1140. A duplicate copy of this form is enclosed. d. <input checked="" type="checkbox"/> The entire content of the foreign application(s), referred to in this application is/are hereby incorporated by reference in this application.					
NOTE: Where an appropriate time limit under 37 C.F.R. 1.494 or 1.495 has not been met, a petition to revive (37 C.F.R. 1.137(a) or (b)) must be filed and granted to restore the application to pending status.					
<div style="display: flex; justify-content: space-between; align-items: flex-end;"> <div> SEND ALL CORRESPONDENCE TO: NIXON & VANDERHYE P.C. 1100 North Glebe Road, 8th Floor Arlington, Virginia 22201 Telephone: (703) 816-4000 </div> <div style="text-align: right;">  SIGNATURE </div> </div>					
B. J. Sadoff NAME					
36,663 REGISTRATION NUMBER					
March 21, 2001 Date					

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Patent Application of

WILSON et al.

Atty. Ref.: 117-347

Serial No. Unknown

Group:

Filed: March 21, 2001

Examiner:

For: TREATMENT OF INFECTION

* * * * *

March 21, 2001

Assistant Commissioner for Patents
Washington, DC 20231

Sir:

PRELIMINARY AMENDMENT

Preliminarily amend the above-identified application as follows.

IN THE SPECIFICATION

Amend the specification as follows.

Insert the attached Sequence Listing in place of the originally-filed copy of the same.

IN THE CLAIMS

Please cancel claims 1-11 without prejudice or disclaimer and add claims 12-19 as follows:

--12. (new) A method of identifying a compound that inhibits the growth of an organism comprising the *ycf 24* gene, the method comprising

- (i) contacting a test compound with the *ycf 24* gene product, and
- (ii) determining whether the test compound inhibits the activity of or binds to the product, any such binding or inhibition being indicative that the compound inhibits the growth of the organism.

13. (new) The method according to claim 12 in which the organism is a malaria parasite.

14. (new) A method of identifying a compound that inhibits the growth of an organism comprising the *ycf 24* gene, the method comprising

- (i) contacting a test compound with a test construct comprising a *ycf 24* promoter operably linked to a coding sequence, and
- (ii) determining whether the test compound inhibits expression driven by the promoter, any such inhibition being indicative that the compound inhibits the growth of the organism.

15. (new) The method according to claim 14 in which the organism is a malaria parasite.

16. (new) A pharmaceutical composition comprising an inhibitor of *ycf 24* gene product expression and/or activity and a pharmaceutically acceptable carrier or diluent.

17. (new) The pharmaceutical composition according to claim 16 wherein the inhibitor is an inhibitor of the growth of a malaria parasite.

18. (new) A method of preventing or treating infection by an organism comprising the *ycf 24* gene in a patient, said method comprising administering to the patient an inhibitor of *ycf 24* gene product expression and/or activity.

19. (new) The method according to claim 18 in which the organism is a malaria parasite.--

REMARKS

The above amendments are made to place the claims in a more traditional format. Claims 12-19 are pending.

WILSON et al.

Serial No. **Unknown**

U.S. National Phase of PCT/GB99/03180

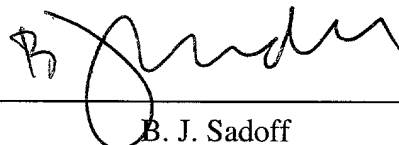
The specification has been amended to include the attached Sequence Listing. The attached paper and computer readable copies of the Sequence Listing are the same. No new matter has been added. A separate Letter stating the same is attached.

An early and favorable Action on the merits is requested.

Respectfully submitted,

NIXON & VANDERHYE P.C.

By: _____



B. J. Sadoff
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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Patent Application of

WILSON et al.

Atty. Ref.: 117-347

Serial No. Unknown

Group:

Filed: March 21, 2001

Examiner:

For: TREATMENT OF INFECTION

* * * * *

March 21, 2001

Assistant Commissioner for Patents
Washington, DC 20231

Sir:

LETTER

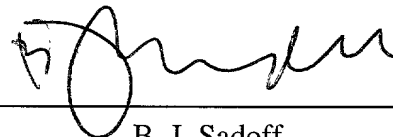
The attached paper and computer readable copies of the Sequence Listing are the same.

No new matter has been added.

Respectfully submitted,

NIXON & VANDERHYE P.C.

By: _____



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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Patent Application of

WILSON et al.

Atty. Ref.: 117-347

Serial No. 09/787,633

Group:

Filed: March 21, 2001

Examiner:

For: TREATMENT OF INFECTION

* * * * *

May 23, 2001

Assistant Commissioner for Patents
Washington, DC 20231

Sir:

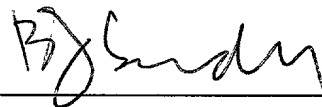
SMALL ENTITY CLAIM

Please accept the Small Entity claim made herein with regard to the above-identified application.

Respectfully submitted,

NIXON & VANDERHYE P.C.

By: _____



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-1-

TREATMENT OF INFECTIONField of the invention

This invention relates to inhibitors of the expression and activity of a gene product and screening methods for the identification of such inhibitors. It further relates to the use of said inhibitors in the treatment of infection by an organism, such as a malaria parasite.

Background to the invention

The malaria parasite and related Apicomplexans are unusual amongst non-photosynthetic organisms in that they possess two forms of organellar DNA, typically a property of plants. One form of organellar DNA is mitochondrial DNA and the other is a 35 kb circle of DNA present in a plastid. The plastid is thought to be of prokaryotic origin. The *ycf 24* gene^{1,2} which is present on the DNA of this plastid corresponds to a well conserved gene which is also found in red algal plastids³ and various bacteria, including *E. coli*.

The plastid genome of Apicomplexans⁴ is distinctive from, but broadly resembles that of the parasitic higher plant *Epifagus virginiana*⁵. It has been suggested that the *raison d'être* for maintenance of vestigial plastid genomes is the small number of open reading frames they carry besides the highly selected set of genes dedicated to the organellar expression system⁶. Unlike Apicomplexans, *Epifagus* does not carry the *ycf 24* gene on its plastid genome, suggesting that either the vestigial organelles have different functions, or a stochastic process determines gene loss or transfer to the nucleus.

The high level of conservation of *ycf 24* (~50% amino acid identity of the putative encoded peptide product with other orthologs) suggests that it is under strong

-2-

selective pressure and likely to have a general function.

Organelle transfection is not available in Apicomplexans. Therefore "knockouts" can only be tested in surrogate organisms.

Summary of the invention

The present inventors have disrupted *ycf 24* in both *Synechocystis* sp. strain PCC6803 and *E. coli*⁷.

Disruption of all the copies of the *ycf 24* gene in these organisms was found to be lethal. Disruption of some of the copies of *ycf 24* was found to give a 'ragged' phenotype in *Synechocystis* and appeared to cause a delay in or interfere in the process of septation during cell division. Thus *ycf 24* is an essential gene and appears to play a role in replication. The present invention is based on the inhibition of the expression or activity of the *ycf 24* product to inhibit the growth of an organism in vivo or ex vivo.

Thus the invention provides an inhibitor of *ycf 24* gene product expression and/or activity for use in a method of treatment of the human or animal body by therapy. Such an inhibitor is useful in particular in the treatment of infection by an organism, such as a malaria parasite. The invention also provides a method of inhibiting the growth of an organism comprising contacting the organism ex vivo with an inhibitor of *ycf 24* gene product expression and/or activity.

Screens may be carried out to identify the inhibitor. Accordingly the invention provides a method of identifying a compound that inhibits the growth of an organism comprising

- (i) contacting a test compound with the *ycf 24* gene product, and
- (ii) determining whether the test compound inhibits the

-3-

activity of or binds to the product, any such binding or inhibition being indicative that the compound inhibits the growth of the organism.

The invention also provides a method of identifying a compound that inhibits the growth of an organism comprising

(i) contacting a test compound with a test construct comprising a *ycf 24* promoter operably linked to a coding sequence,

(ii) determining whether the test compound inhibits expression driven by the promoter, any such inhibition being indicative that the compound inhibits the growth of the organism.

The invention further provides a pharmaceutical composition comprising an inhibitor of the invention and a pharmaceutically acceptable carrier or diluent.

As noted above the inhibitor may be used to treat an infection by an organism and therefore the invention provides a method of treating a host suffering from an infection by an organism, which method comprises administering to the host a therapeutically effective amount of an inhibitor of *ycf 24* product expression and/or activity.

Brief description of the drawings

The invention is illustrated by the accompanying drawings in which:

Fig.1 shows phenotypic characters of wild type (WT), ragged transformant (R) and smooth mutant forms (S) of *Synechocystis* sp. : a) Colony shape and size; an enlargement is inserted in R, b) Scanning electron micrographs of whole cells - bar = 10µm, c) Cross-sections showing plaques - arrow, d) Whole cells with and without 4',6-diamindino-2-phenylindole (DAPI)-staining. Arrows point to mis-segregation.

-4-

Fig.2 shows a Southern blot of DNA from *Synechocystis* sp., restricted with *Hind* III and hybridized with wild type (WT) *ycf* 24. The sizes expected for restriction fragments from WT (lane 4) and clones of heteroplasmic transformants (lanes 1-3) are given in the diagram below.

Fig.3 shows an effect of over-expression of *ycf* 24 in *E.coli*. Fig. 3a shows nucleoids stained with DAPI in an uninduced culture of *E.coli* and Fig. 3b shows nucleoids stained with DAPI after induction of over-expression.

Detailed description of the invention

The inhibitor can be used to inhibit the growth of a unicellular or multicellular organism in vivo or ex vivo. The organism may be a prokaryote or a eukaryote. The organism comprises the *ycf* 24 gene. In the case of a eukaryote the nuclear genome of the organism may contain the *ycf* 24 gene or the organism may contain an organelle which contains DNA comprising the *ycf* 24 gene. Such an organelle is generally of prokaryotic origin, for example algal origin, and may be a plastid.

The organism may be a protozoa, such as an Apicomplexan, for example of clade plasmodium, piroplasm, sarcocystan or coccidium. The organism may be *Plasmodium falciparum*. The organism may be an alga. The organism may be a bacterium, such as a cyanobacterium, mycobacterium or *E. coli*. The organism may be free-living or a pathogen of humans and/or animals. It may be an intracellular or extracellular pathogen.

The *ycf* 24 gene product is generally one which can be expressed from the coding region of:

(a) the polynucleotide sequence of SEQ ID NO: 1, 2 or 3

-5-

or a fragment thereof; or

(b) polynucleotides which can selectively hybridise to the coding region of (a) of a fragment thereof; or

(c) polynucleotides which, but for the degeneracy of the genetic code, would hybridise to (a) or (b).

The polynucleotides of (b) may selectively hybridise under conditions of medium to high stringency, for example 0.03M NaCl and 0.03M sodium citrate at from 50 to 60 degrees centigrade.

Thus the polynucleotide from which the *ycf* 24 product is expressed generally has at least 70%, preferably at least 95, 97 or 99% sequence identity to the coding sequence of SEQ ID NO: 1, 2, or 3 over a region of at least 20, preferably at least 30, for instance at least 46, 60 or 100 or more contiguous nucleotides.

The *ycf* 24 product generally has at least 50% sequence identity to the amino acid sequences shown is SEQ ID NO: 1, 2, or 3, preferably at least 80 or 90% and more preferably at least 95, 97 or 99% over a region of at least 20, preferably at least 30, for instance at least 46, 60, 100 or more contiguous amino acids. The term *ycf* 24 product includes fragments of the amino acid sequences of SEQ ID NO: 1, 2 or 3 or of the homologous sequences discussed above. Suitable fragments will be at least 5, 10, 20 or 40 amino acids in size, for example at least 60 or 100 amino acids in size. Generally the *ycf* 24 product has *ycf* 24 product activity, and would generally be able to complement the activity of a naturally occurring *ycf* 24 product.

Any suitable inhibitor of *ycf* 24 product activity may be used in the present invention. The inhibitor may bind in a reversible or irreversible manner. An irreversible inhibitor would dissociate very slowly from the *ycf* 24 product because it would be very tightly

-6-

bound, either covalently or non-covalently. Reversible binding, in contrast with irreversible binding, is characterised by a rapid dissociation of the *ycf 24* product-inhibitor complex.

5 As noted above the invention provides a method of identifying a compound that inhibits the activity of *ycf 24* product. *ycf 24* product for use in this method can be obtained, for example, recombinantly by any method known in the art. The nucleotide sequences of the *ycf 24*
10 genes of *Plasmodium falciparum*, *Synechocystis* Sp strain pCC6803 and *E.coli* are provided herein as SEQ ID NOs 1, 2 and 3 respectively. The *ycf 24* genes of other species may be identified by searching in databases for genes homologous to SEQ ID NO: 1, 2 or 3. Nucleotides with the
15 same sequence as genes identified in a database can be made and expressed using routine methods which are known in the art. Alternatively *ycf 24* genes can be identified by probing genomic or cDNA libraries with nucleotides comprising the *ycf 24* gene or a fragment of the *ycf 24*
20 gene and then isolating and sequencing the nucleotides identified by the probe.

Any suitable format may be used for identifying the inhibitor of the activity. In the method the *ycf 24* product is generally in a suitable buffer, which includes
25 any suitable biological buffer that can provide buffering capability at a pH conducive to the reaction requirements of the *ycf 24* product. The *ycf 24* product may be in conditions which are similar to the physiological conditions in which it occurs naturally.

30 In the method the *ycf 24* product may be in inside a cell or outside a cell. The cell may be the cell in which the *ycf 24* product naturally occurs, or a cell in which the *ycf 24* product is expressed recombinantly. The cell may be treated with agents which permeabilise the cell

-7-

surface allowing test substances to enter the cell more readily. The *ycf* 24 product used in the method may be in the form of an extract from such cells.

5 The method may determine whether a test compound is able to bind the *ycf* 24 product, such as in a specific manner. This method may comprise allowing the *ycf* 24 product to bind to a test compound and measuring the amount of binding that occurs, such as by measuring the amount of *ycf* 24 bound to the compound. The amount of
10 binding may be determined by measuring a characteristic of the *ycf* 24 product that changes upon binding, such as spectroscopic changes.

15 The assay format may be a 'band shift' system. This involves determining whether a test compound advances or retards the *ycf* 24 product on gel electrophoresis relative to the *ycf* 24 product in the absence of the compound.

20 The method may be a competitive binding assay. This determines whether the test compound is able to inhibit the binding of the *ycf* 24 product to a substance which is known to bind to the *ycf* 24 product, such as an antibody specific for product. This method allows test compounds to be identified which are 'analogues' of *ycf* 24 product.

25 The competitive binding system may comprise
(i) incubating the *ycf* 24 product with a test compound and a labelled reference compound that is known to bind the product;
(ii) determining the amount of the labelled reference compound that is bound to the product; and
30 (iii) comparing the amount of bound labelled reference compound determined in step (ii) with the amount of said compound that binds to the product in the absence of the test compound;
wherein any reduction in the binding of the labelled

-8-

reference compound in the presence of the test compound compared to the binding in the absence of the test compound shows that the test compound is competing with the reference compound for binding to the product. This indicates that the test compound could be an inhibitor of *ycf 24* product activity.

The amount of the labelled reference compound bound to the *ycf 24* product may be measured directly or indirectly. A direct measurement may be carried out by removing assay mixture containing the unbound labelled reference compound and measuring the amount of label that is in the product fraction. Alternatively, the amount of labelled reference compound bound to the product could be determined indirectly by measuring the amount of label remaining in the assay solution after removal of the product fraction, which will be inversely related to the amount that has bound to the product.

In a competitive binding assay system, the *ycf 24* product may be immobilised on a solid support or may be in solution. The use of immobilised product has the advantage that, after the binding reaction is complete, the product/labelled reference compound complex may be separated from the labelled reference compound that remains in solution by simply removing the solution away from the solid support. If, on the other hand, the product is not immobilised during the assay but rather is in solution, then it will generally be necessary to devise a means for separating the product/labelled reference compound complex from the uncomplexed reference compound before measuring the amount of label. Such separation could be achieved, for example, by precipitating the product using an antibody to the product or by using a non-specific precipitation technique.

Suitable labels for use in the assay systems

-9-

according the invention are well-known in the art. The reference compound which is know to bind the *ycf* 24 product may be an antibody.

5 An antibody to *ycf* 24 product may be produced by raising antibody in a host animal against the whole *ycf* 24 product or an antigenic epitope thereof (hereinafter "the immunogen"). Methods of producing monoclonal and polyclonal antibodies are well-known. A method for producing a polyclonal antibody comprises
10 immunising a suitable host animal, for example an experimental animal, with the immunogen and isolating immunoglobulins from the serum. The animal may therefore be inoculated with the immunogen, blood subsequently removed from the animal and the IgG fraction purified. A method for producing a monoclonal antibody comprises
15 immortalising cells which produce the desired antibody. Hybridoma cells may be produced by fusing spleen cells from an inoculated experimental animal with tumour cells (Kohler and Milstein, Nature 256, 495-497, 1975).

20 An immortalized cell producing the desired antibody may be selected by a conventional procedure. The hybridomas may be grown in culture or injected intraperitoneally for formation of ascites fluid or into the blood stream of an allogenic host or
25 immunocompromised host. Human antibody may be prepared by *in vitro* immunisation of human lymphocytes, followed by transformation of the lymphocytes with Epstein-Barr virus.

30 For the production of both monoclonal and polyclonal antibodies, the experimental animal is suitably a goat, rabbit, rat or mouse. If desired, the immunogen may be administered as a conjugate in which the immunogen is coupled, for example via a side chain of one of the amino acid residues, to a suitable carrier. The carrier
35 molecule is typically a physiologically acceptable

-10-

carrier. The antibody obtained may be isolated and, if desired, purified.

Any suitable inhibitor of expression of *ycf 24* product may be employed in the present invention.

5 Preferably the inhibitor is a specific inhibitor of transcription from the *ycf 24* gene, and does not inhibit transcription from other genes.

10 The inhibitor may inhibit transcription or translation of *ycf 24* product. The inhibitor may bind to the *ycf 24* gene either 5' to the coding sequence, to the coding sequence or 3' to the coding sequence. Thus the inhibitor may bind to the *ycf 24* promoter, and inhibit the initiation of transcription. The inhibitor may bind and inhibit the action of a protein which is required for transcription from the *ycf 24* gene. The inhibitor may bind to the untranslated or translated regions of the *ycf 24* mRNA. This could prevent the initiation of translation. Alternatively the inhibitor could bind to a protein which associates with the untranslated region and prevent the protein associating with the untranslated region.

Any suitable assay format may be used for identifying an inhibitor of *ycf 24* expression. The invention provides a method for identifying an inhibitor of *ycf 24* expression comprising:

(i) contacting a compound with a nucleic acid comprising a *ycf 24* promoter operably linked to a coding sequence; and
(ii) determining whether the compound inhibits expression driven by the promoter.

30 Generally this method is carried out in conditions which in the absence of the test compound lead to expression of the coding sequence from the nucleic acid. The nucleic acid may also comprise other untranscribed or untranslated regions of the *ycf 24* gene. The coding

-11-

sequence typically encodes a protein that is able to act as a reporter of expression. The assay may be carried out in a cell which harbours the nucleic acid. The substance may be tested with any other known promoter to test the possibility that the test substance is a general inhibitor of gene expression.

Any reporter polypeptide may be used, for example GUS or GFP. GUS is assayed by measuring the hydrolysis of a suitable substrate, for example 5-bromo-4-chloro-3-indolyl- β -D-glucuronic acid (X-gluc) or 4-methylumbelliferyl- β -glucuronide (MUG). The hydrolysis of MUG yields a product which can be measured fluorometrically. GFP is quantified by measuring fluorescence at 590nm after excitation at 494nm. These methods are well known to those skilled in the art.

Alternatively the coding sequence may be the *ycf 24* coding sequence itself, or a fragment of this sequence. The expression of the *ycf 24* product may be measured by for example, Northern/RNA blotting, Western/antibody blotting, RNA in situ hybridization or immunolocalisation. Thus the method for identifying an inhibitor of expression of *ycf 24* product may be performed on a cell which naturally harbours the *ycf 24* gene, or an extract from such a cell.

Suitable candidate substances which may be inhibitors and which can be tested in the methods discussed above include antibody products (for example, monoclonal and polyclonal antibodies, single chain antibodies, chimeric antibodies and CDR-grafted antibodies) which are specific for *ycf 24* product. Furthermore, combinatorial libraries, defined chemical identities, peptide and peptide mimetics, oligonucleotides and natural product libraries, such as display libraries (e.g. phage display libraries) may also be tested. The candidate substances may be chemical

-12-

compounds. Batches of the candidate substances may be used in an initial screen of, for example, ten substances per reaction, and the substances of batches which show inhibition tested individually.

5

Antisense inhibitors

As noted above the expression of *ycf 24* in a cell may be reduced by the presence in that cell of a compound which can bind to the *ycf 24* mRNA. Therefore a polynucleotide which is capable of hybridizing to *ycf 24* mRNA can
10 constitute an appropriate inhibitor of *ycf 24* expression.

The polynucleotide may be antisense to the *ycf 24* mRNA. Such a polynucleotide may be capable of hybridising to *ycf 24* mRNA and may thus inhibit the expression of *ycf 24* product by interfering with one or
15 more aspects of *ycf 24* mRNA metabolism including processing, translation and metabolic turnover.

The antisense polynucleotide may be RNA. This may hybridise to the *ycf 24* mRNA to form an RNA-RNA duplex which may cause direct inhibition of translation and/or the destabilization of the target message, for example, rendering susceptibility to nucleases.
20

Such a polynucleotide may hybridize to all or part of the *ycf 24* mRNA. Typically the antisense
25 polynucleotide hybridizes to the ribosome binding region or the coding region of the *ycf 24* mRNA. The polynucleotide may be complementary to all of or a region of the *ycf 24* mRNA. For example, the polynucleotide may be the exact complement of all or a part of *ycf 24* mRNA.
30 However, absolute complementarity is not required and polynucleotides which have sufficient complementarity to form a duplex having a melting temperature of greater than 20°C, 30°C or 40°C under physiological conditions are particularly suitable for use in the present invention.

-13-

The polynucleotide may be a polynucleotide which hybridises to the *ycf 24* mRNA under conditions of medium to high stringency such as 0.03M sodium chloride and 0.03M sodium citrate at from about 50 to about 60 degrees centigrade.

It is preferred that the polynucleotide hybridizes to all or part of the region of the *ycf 24* mRNA corresponding to the coding sequence defined by nucleotides 26 to 1435 of SEQ ID NO:1. However, the polynucleotide may hybridise to all or part of the 5'- or 3'-untranslated region of the mRNA. The polynucleotide will typically be from 6 to 40 nucleotides in length. Preferably it will be from 12 to 20 nucleotides in length. The polynucleotides may be at least 40, for example at least 60 or at least 80, nucleotides in length and up to 100, 200, 300, 400, 500, 600 or 700 nucleotides in length or even up to a few nucleotides, such as five or ten nucleotides, shorter than SEQ ID NO: 1, 2 or 3.

When the polynucleotide is an antisense RNA it may be expressed in a cell from a recombinant replicable vector. Such a replicable vector comprises a polynucleotide which when transcribed gives rise to antisense RNA.

Thus the antisense polynucleotide may be provided by delivering such a vector to the cell and allowing transcription from the vector to occur. Such a vector is also understood to an inhibitor of *ycf 24* expression in this specification. Generally on the vector the polynucleotide giving rise to the antisense RNA is operably linked to a control sequence which is capable of providing for the transcription of the polynucleotide giving rise to the antisense RNA. The term "operably linked" refers to a juxtaposition wherein the components described are in a relationship permitting them to function in their intended manner. A control sequence

-14-

"operably linked" to a sequence giving rise to an antisense RNA is ligated in such a way that transcription of the sequence is achieved under conditions compatible with the control sequences.

5 The vector may be for example, a plasmid or virus vector provided with an origin of replication, optionally a promoter for transcription to occur and optionally a regulator of the promoter. The vector may contain one or more selectable marker genes, for example an ampicillin
10 resistance gene in the case of bacterial plasmid or a neomycin resistance gene for a mammalian vector. Vectors may be used in vitro, for example for the production of antisense RNA or used to transfect or transform a host cell. The term 'host cell' refers either to the organism
15 whose growth it is desired to inhibit, or a human or animal cell which can be infected by the organism. Thus the host cell may be an infected cell. The vector may also be adapted to be used in vivo, for example in a method of gene therapy.

20 Promoters/enhancers and other expression regulation signals may be selected to be compatible with the host cell for which the expression vector is designed. For example, mammalian promoters, such as b-actin promoters, may be used. Tissue-specific promoters, in particular
25 neuronal cell specific promoters (for example the tyrosine hydroxylase (TH), L7, or neuron specific enolase (NSE) promoters), are especially preferred. Viral promoters may also be used, for example the Moloney murine leukaemia virus long terminal repeat (MMLV LTR),
30 the rous sarcoma virus (RSV) LTR promoter, the SV40 promoter, the human cytomegalovirus (CMV) IE promoter, herpes simplex virus promoters or adenovirus promoters. All these promoters are readily available in the art. Preferred promoters are tissue specific promoters such as
35 the casein gene promoter.

-15-

The vector may further include sequences flanking the polynucleotide giving rise to antisense RNA which comprise sequences homologous to eukaryotic genomic sequences, preferably mammalian genomic sequences, or viral genomic sequences. This will allow the introduction of the polynucleotides of the invention into the genome of eukaryotic cells or viruses by homologous recombination. In particular, a plasmid vector comprising the expression cassette flanked by viral sequences, for example HSV1 or HSV2 sequences, can be used to prepare a viral vector, for example an HSV vector, suitable for delivering the polynucleotides of the invention to a mammalian cell. Other examples of suitable viral vectors include retroviruses, including lentiviruses, adenoviruses and adeno-associated viruses. Gene transfer techniques using these viruses are known to those skilled in the art. Retrovirus vectors for example may be used to stably integrate the polynucleotide giving rise to the antisense RNA into the host genome. Replication-defective adenovirus vectors by contrast remain episomal and therefore allow transient expression.

Antisense polynucleotides may be chemically modified. This may enhance their resistance to nucleases and may enhance their ability to enter cells. For example, phosphorothioate oligonucleotides may be used. Other deoxynucleotide analogs include methylphosphonates, phosphoramidates, phosphorodithioates, N3'P5'-phosphoramidates and oligoribonucleotide phosphorothioates and their 2'-O-alkyl analogs and 2'-O-methylribonucleotide methylphosphonates.

Alternatively mixed backbone oligonucleotides (MBOs) may be used. MBOs contain segments of phosphothioate oligodeoxynucleotides and appropriately placed segments of modified oligodeoxy- or oligoribonucleotides. MBOs have segments of phosphorothioate linkages and other segments of

-16-

other modified oligonucleotides, such as methylphosphonate, which is non-ionic, and very resistant to nucleases or 2'-O-alkyloligoribonucleotides.

An inhibitor of *ycf 24* expression and/or activity is one which produces a measurable reduction in *ycf 24* expression and/or activity in the methods described above. Preferred substances are those which inhibit *ycf 24* expression and/or activity by at least 10%, at least 20%, at least 30%, at least 40% at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95% or at least 99% at a concentration of the inhibitor of $1\mu\text{g ml}^{-1}$, $10\mu\text{g ml}^{-1}$, $100\mu\text{g ml}^{-1}$, $500\mu\text{g ml}^{-1}$, 1mg ml^{-1} , 10mg ml^{-1} or 100mg ml^{-1} . The percentage inhibition represents the percentage decrease in expression/activity in a comparison of assays in the presence and absence of the test substance. Any combination of the above mentioned degrees of percentage inhibition and concentration of inhibitor may be used to define an inhibitor of the invention, with greater inhibition at lower concentrations being preferred.

When used in vivo or in vitro the inhibitor inhibits the growth of the organism. Generally the inhibitor will decrease the rate of cell division of the organism. Generally the inhibitor will decrease the number of divisions that occur in a unit of time by at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or 100% at a concentration of inhibitor of $1\mu\text{g ml}^{-1}$, $10\mu\text{g ml}^{-1}$, $100\mu\text{g ml}^{-1}$, $500\mu\text{g ml}^{-1}$, 1mg ml^{-1} , 10mg ml^{-1} or 100mg ml^{-1} .

If the organism is one in which the *ycf 24* gene is present in an organelle then the inhibitor generally decreases the rate of division of that organelle. The inhibitor may also disrupt cell or organelle division in other ways, such as causing abnormal septation or mis-segregation of DNA. The inhibitor may also disrupt the structure of the organism, such as giving the organism a 'ragged' appearance. The inhibitor may cause the death of

-17-

the organism.

Candidate substances which show activity in assays such as those described herein can then be tested on the organism in question. The inhibitor may or may not be toxic towards humans or animals. The inhibitor may thus be used as an antibiotic.

The inhibitors of *ycf 24* expression and/or activity or the vectors described above may be used to treat infections by an organism and, in particular, malaria. The condition of a patient suffering from an infection can therefore be improved by administration of such an inhibitor. A therapeutically effective amount of such an inhibitor may be given to a human patient in need thereof.

The formulation of inhibitor or vector for use in preventing or treating infection by an organism will depend upon factors such as the nature of the substance identified, whether a pharmaceutical or veterinary use is intended, etc. In order to be administered to a patient, the compound will be provided in the form of a pharmaceutical composition containing the inhibitor or vector and a pharmaceutically acceptable carrier or diluent. Suitable carriers and diluents include isotonic saline solutions, for example phosphate-buffered saline. Typical oral dosage compositions include tablets, capsules, liquid solutions and liquid suspensions. For example it may be formulated for parenteral, intravenous, intramuscular, subcutaneous, transdermal or oral administration.

The dose of inhibitor may be determined according to various parameters, especially according to the substance used; the age, weight and condition of the patient to be treated; the route of administration; and the required regimen. A physician will be able to determine the required route of administration and dosage for any particular patient. A suitable dose may however be from

-18-

0.1 to 100 mg/kg body weight such as 1 to 40 mg/kg body weight, for example, to be taken from 1 to 3 times daily.

The oral route is preferred because this is generally the most convenient route for a patient to take regular doses of the compound without the assistance of a physician. However, the inhibitor or vector may be administered by direct injection into the site to be treated.

The inhibitor or vector may be used in either the treatment of an existing infection by the organism or in the prevention of such an infection from occurring in the first place.

The vectors giving rise to antisense RNA of the invention may be administered directly as a naked nucleic acid construct. Uptake of naked nucleic acid constructs by mammalian cells is enhanced by several known transfection techniques, for example those including the use of transfection agents. Example of these agents include cationic agents (for example calcium phosphate and DEAE-dextran) and lipofectants (for example lipofectamTM and transfectamTM). These agents may also enhance the uptake of the constructs by the organism.

The pharmaceutical composition may be formulated to aid delivery of the inhibitor or vector to the organism whose growth is to be inhibited. In particular it may be formulated to deliver the inhibitor or vector to the cell type which is infected or to the compartment in the human or animal cell which is infected by the organism. It may be formulated to deliver the inhibitor or vector to the organelle in the organism where *ycf 24* product or mRNA is present.

When the pharmaceutical composition comprises viral vector for gene therapy it may be administered in such a way that the vector is incorporated into cells at an appropriate area. The amount of virus administered is in

-19-

the range of from 10^4 to 10^8 pfu, preferably from 10^5 to 10^7 pfu, more preferably about 10^6 pfu for herpes viral vectors and from 10^6 to 10^{10} pfu, preferably from 10^7 to 10^9 pfu, more preferably about 10^8 pfu for adenoviral vectors. When injected, typically 1-2 ml of virus in a pharmaceutically acceptable suitable carrier or diluent is administered. When the polynucleotide is administered as a naked nucleic acid, the amount of nucleic acid administered is typically in the range of from 1 μ g to 10 mg.

Where the polynucleotide giving rise to the antisense RNA is under the control of an inducible promoter, it may only be necessary to induce gene expression for the duration of the treatment. Once the condition has been treated, the inducer is removed and expression of the polypeptide of the invention ceases. This will clearly have clinical advantages. Such a system may, for example, involve administering the antibiotic tetracycline, to activate gene expression via its effect on the tet repressor/VP16 fusion protein.

The use of tissue-specific promoters will be of assistance in the treatment of disease using the vectors of the invention. It will be advantageous to be able express inhibitors of *ycf 24* in only the relevant affected cell types, especially where such inhibitors are toxic when expressed in other cell types.

When the inhibitor is used to inhibit the growth of an organism ex vivo the organism may or may not be a pathogen of humans or animals. The inhibitor may be used to treat land, water or food containing the organism.

The following Examples illustrate the invention:

Example 1

Gene disruption of *ycf 24*, transformation and selection

A central segment of the wild type gene (*slr0074*) from *Synechocystis* sp., strain pCC6803, (delineated by

-20-

two *Hind*III sites ~1.0 kb apart) was amplified from genomic DNA by PCR using two oligonucleotide primers based on the known sequence (Accession no. S76598): 5' CTC CGC CCC TAA GCA AAG TAA GAA A and 5' TGA TCC TCG CCA ATT TTG GAA GTG GA. The amplified fragment was restricted with *Hind*III and cloned into pUC9 to give pUC9/slr. The sequence of both ends of the insert was confirmed in recombinants grown in Epicurian SURE cells. To disrupt the gene, a kanamycin resistance gene (k^r) isolated from a modified bluescript vector (pBSHdSpl, was blunt-ended with the Klenow fragment and cloned into a unique *Xba*I site in the *ycf* 24 insert to give pUC9/slr/ k^r . Recombinants in XL1-blue MRF' cells were selected using kanamycin. Disruption of the *ycf* 24 insert was confirmed by sequencing the junction regions.

Wild type (WT) cells of *Synechocystis* sp. strain PCC 6803 were transfected by incubation with light at 30°C for 4 hr with vector DNA carrying the disrupted *ycf* 24 gene (1-5 µg in 10 µl water). Selection and growth were carried out on BG11 plates under photo-autotrophic conditions at 30°C. Several rounds of replication were allowed over 3 days before recombinants were selected by overlaying with kanamycin (50 µg/ml). This killed most of the cells but transformants emerged after a delay. Every 10-14 days thereafter, transformants were picked and re-plated to allow segregation of mutant cells. For growth comparisons in the absence of selection, WT cells were plated at the same time on BG11 (their growth was inhibited totally by kanamycin at 50 µg/ml).

Example 2

Gene Disruption in *E. coli*

Disruption of *ycf* 24 in *E. coli* was carried out along similar lines, but in this case disruption was with the *aadA* gene for streptomycin resistance (S^r). *ycf* 24 from *E.*

-21-

coli was amplified using primers based on the accession no. D90811 with added terminal restriction sites: 5' GAG CTC GGA ATT CGC ATG TGG CTG TGG CGA AAG and 3' GAG CTC GGG ATC CTT ATC CGA CGC TGT GTT CAA G. The *E.coli aadA* gene was introduced at a *BsgI* restriction site near the centre of *ycf 24* and cloned into the bluescript vector pBSKS⁺. The construct was linearized for transformation and to allow homologous recombination in *E. coli* LE392 host cells. The linearized recombinant plasmid pBSKS⁺/*ycf24*/*S^r* was transfected into *E. coli* by heat shock but no homologous recombinants were found from the primary normal size colonies; a few antibiotic resistant colonies were recovered that carried circular episomes likely to have arisen from re-ligation of the construct. After 2 days incubation at 37°C small colonies were observed. Cells from these were found to be elongated (at least x3 length of host cells) but they survived for only a few generations. PCR from these colonies to determine if a "knock-out" had occurred was unsuccessful.

Example 3

Characterising the *Synechocystis* transformants

WT *Synechocystis* and transformants carrying an homologous form of *ycf 24* interrupted by a kanamycin resistance gene (*k^r*) were plated on BG-11 agar. The WT cells formed colonies visible to the naked eye within 4 days, whereas it took 2 weeks for transformants to appear under selection with kanamycin (50 µg ml⁻¹). Colonies of transformants were small and "ragged" in outline in contrast to the larger, smooth-contoured WT colonies (Fig. 1a). DNA extracted from cloned transformants was hybridized in Southern blots with WT *ycf 24* and found to be heteroplasmic, the copy numbers of the disrupted and WT forms of *ycf 24* being approximately equal as judged from

-22-

the intensity of the hybridization signals (Fig. 2). Relaxation of kanamycin selection on heteroplasmic transformants resulted in vigorously growing, smooth-contoured colonies - "revertants", confirming that the heteroplasmic state was growth-limiting. These observations were confirmed by additional transformations, including one using disruption with a spectinomycin resistance gene instead of kanamycin. Transformants homoplasmic for the disrupted form of *ycf 24* were never recovered, even after prolonged selection with kanamycin at 50 or 100 $\mu\text{g ml}^{-1}$ and homoplasmy is inferred to be lethal.

Longer-term selection of "ragged" transformants on kanamycin also gave rise to rapidly growing colonies with smooth contours. However, hybridization on Southern blots showed these remained heteroplasmic for the disrupted gene and we refer to these as "smooth" suppressors. DNA from cloned "smooth" suppressors was used to transform both WT and "ragged" heteroplasmic cells. In both cases, kanamycin selection yielded mostly "ragged" colonies but small numbers (<1%) of smooth, rapidly growing colonies were found, unlike in the original transfections.

The cells were examined by electron microscopy. Pelleted cells were pre-fixed in 0.5% neutral glutaraldehyde in 100mM cacodylate buffer, pH 7.0 for 30 min at room temperature. After gentle centrifugation, the pelleted cells were fixed in 2-3% glutaraldehyde buffer and thin sections were prepared using standard procedures. For scanning electron microscopy (SEM), cells were spread on glass slips coated in poly-L-lysine (100 $\mu\text{g ml}^{-1}$) and fixed in 2% glutaraldehyde. Post fixation samples were washed in ethanol and rehydrated by immersion in hexamethyl-disilizane. After overnight dessication, samples were sputter-coated with 20 nm gold (for sample conductivity) and examined by SEM.

Electron micrographic sections showed ~80% of cells

-23-

from "ragged" colonies (>100 cells examined) had electron-translucent plaques surrounding the phycobilisomes in the thylacoid membranes (Fig. 1c). The number of cells containing plaques fell to ~45% upon relaxation of kanamycin selection pressure, coincident with reversion to normal colony morphology. Because similar plaques were found in ~4% WT cells, we suggest they indicate "sick" cells rather than a direct phenotypic change following disruption of *ycf 24*.

By contrast, examination of the gross morphology of cells from "ragged" colonies by scanning electron microscopy showed a significant accumulation of cells (~35%) in the last stages of cell division ($p = <0.001$), suggesting a defect in cytokinesis (Table 1 and Fig.1b). Relaxation of kanamycin selection again saw the proportion of dividing cells in the transformed cell population return to one resembling WT. A high proportion (~60%) of the "smooth" suppressor cells also accumulated at a late stage of cytokinesis, with occasional small chains of incompletely separated cells. Smooth suppressors still contained electron-opaque plaques and electron micrographic sections confirmed that suppression increased the frequency of aberrant septation in dividing cells.

Table 1

Cell type ^{*1}	% Fission			P (X^2) ^{*3}
	Early stage	Late stage	Total	
WT	10	9 ^{±1}	19	-
Ragged	21 ^{*6}	34 ^{±1}	55	<0.001
Revertant ^{*2}	8	4	12	0.9
Smooth	20	41	61	<0.001

^{*1} Nos. cells counted = >100/group (mean data from 2 exp.)

-24-

*² Ragged transformants grown without kanamycin selection

*³ Chi-square comparison with WT

DNA was stained with 4',6-diamindino-2-phenylindole (DAPI) at 0.01µg ml⁻¹, following a protocol similar to that described¹⁵. When stained with DAPI, the nucleoids of WT cells were highly uniform in appearance (Fig.1d). Most WT cells (~82%) were not in division and only a few dead cells (unstained) were present. DAPI-staining of cells from "ragged colonies" was much more variable, ranging from cells that looked like WT to a large number (~50%) of dead or dying cells that stained less intensely. The latter included >50% of those in division. Although nucleoid segregation had occurred in many of these cells, in ~40% the DNA had mis-segregated (ie. one daughter cell was without DNA or carried less DNA as judged by staining intensity); altered autofluorescence under UV light indicated these were dying cells. DAPI-stained "smooth" suppressants also were distinctive. About half of the cells (both dividing and non-dividing) carried normal amounts of DNA but occasional cells either had no DNA (~1% dead), had undergone asymmetric DNA segregation (~1%), or had about twice as much DNA (~1%) as WT. The cells in this last group were larger than the rest (Fig. 1d). Coulter counter analysis of the size distribution of the different populations (data not shown) confirmed that the numbers of small and large cells had substantially increased in both the "ragged" and smooth isolates compared with WT. The amount of DNA per cell corresponded generally with cell size, excepting dead cells.

Example 4

Over-expression of ycf 24 in *E.coli*

The effect of over-expressing ycf 24 in *E.coli* was examined using the expresson vector pMAL-c2.

-25-

Gene amplification was carried out with primers designed to give a 5' Eco RI site and a 3' Pst I site, and the product was cloned into the pMAL-c2 expression vector (New England Biolabs). Cultures of transfectants growing exponentially in Terrific broth containing ampicillin (50 mg ml⁻¹) were induced with 0.3-0.5 mM IPTG. Transformants expressing fusion proteins after induction at 37°C were detected by Western blotting and by immunofluorescence using antibody to the maltose binding protein tag of the fusion protein.

Filamentation was marked in such transformants.

Immunofluorescence showed a striking "zebra" pattern of the fusion protein which was mainly sequestered on either side of the nucleoids of the bacteria.

After prolonged over-expression, the bands of fusion protein appeared at other sites along the length of the extended cells where there were no nucleoids. This correlated with aberrant partition of the nucleoids. We found the nucleoids divided but did not separate, forming a clump that distorted the cell (see Fig. 3). We also noted that the first division of the nucleoids was parallel and not transverse to the cell length so that nucleoids lay side by side rather than being displaced longitudinally as is usual. Deletion analysis of the *ycf 24* gene showed that the "zebra" pattern due to deposition of the fusion protein depends on a sequence lying between amino acids 81 and 161 at the N-terminus of the protein. Secondary structure analysis of *ycf 24* indicates it probably has several domains, two of which lie in the highly conserved C-terminal half. One domain resembles a serine protease but lacks the catalytic site. Another, at the C-terminus itself, is folded like complement factor 5a.

Examination of the cells over-expressing *ycf 24* showed they contained only slightly depressed levels of the essential cell division protein FtsZ, yet no septal rings

-26-

were formed (unlike in some other nucleoid partition mutants of *E.coli*). Thus, over-expression of *ycf 24* seems to disturb formation of the septal ring complex required for cell division.

5 Our studies with *E.coli* and *Synechocystis* concur in showing that modulating the amount of *ycf 24* affects cell division. Work is in progress to ascertain the function of the protein by determining its interactions with other cellular constituents. This includes a detailed analysis
10 of the sequence required for cellular localization which might be susceptible to disruption by an antisense approach.

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SEQUENCE LISTING

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-2-

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1344
 1392
 1440
 1488
 1527
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-28-

Claims

1. An inhibitor of ycf 24 gene product expression and/or activity for use in a method of treatment of infection of the human or animal body by an organism comprising said gene.

2. An inhibitor according to claim 1 for use in a method of treatment of malaria.

10

3. A method of inhibiting the growth of an organism comprising the ycf 24 gene, which method comprises contacting the organism ex vivo with an inhibitor of expression and/or activity of the gene product.

15

4. A method of identifying a compound that inhibits the growth of an organism comprising the ycf 24 gene, the method comprising

20

(i) contacting a test compound with the ycf 24 gene product, and

(ii) determining whether the test compound inhibits the activity of or binds to the product, any such binding or inhibition being indicative that the compound inhibits the growth of the organism.

25

5. A method of identifying a compound that inhibits the growth of an organism comprising the ycf 24 gene, the method comprising

(i) contacting a test compound with a test construct comprising a ycf 24 promoter operably linked to a coding sequence,

(ii) determining whether the test compound inhibits expression driven by the promoter, any such inhibition being indicative that the compound inhibits the growth of the organism.

35

-29-

6. A method according to claim 4 or 5 in which the organism is a malaria parasite.

7. A method of inhibiting the growth of an organism comprising the *ycf 24* gene, said method comprising contacting the organism *ex vivo* with a compound identified by the method of any one of claims 4 to 6.

8. Use of a compound identified by the method of any one of claims 4 to 6 for the manufacture of a medicament for the treatment of an infection by an organism comprising the *ycf 24* gene.

9. A pharmaceutical composition comprising an inhibitor of *ycf 24* gene product expression and/or activity or a compound identified by the method of any one of claims 4 to 6 and a pharmaceutically acceptable carrier or diluent.

10. A method of preventing or treating infection by a unicellular organism comprising the *ycf 24* gene in a patient, said method comprising administering to the patient an inhibitor of *ycf 24* gene product expression and/or activity or a compound identified by the method of any one of claims 4 to 6.

11. A method according to claim 10 in which the organism is a malaria parasite.

Fig.1a.

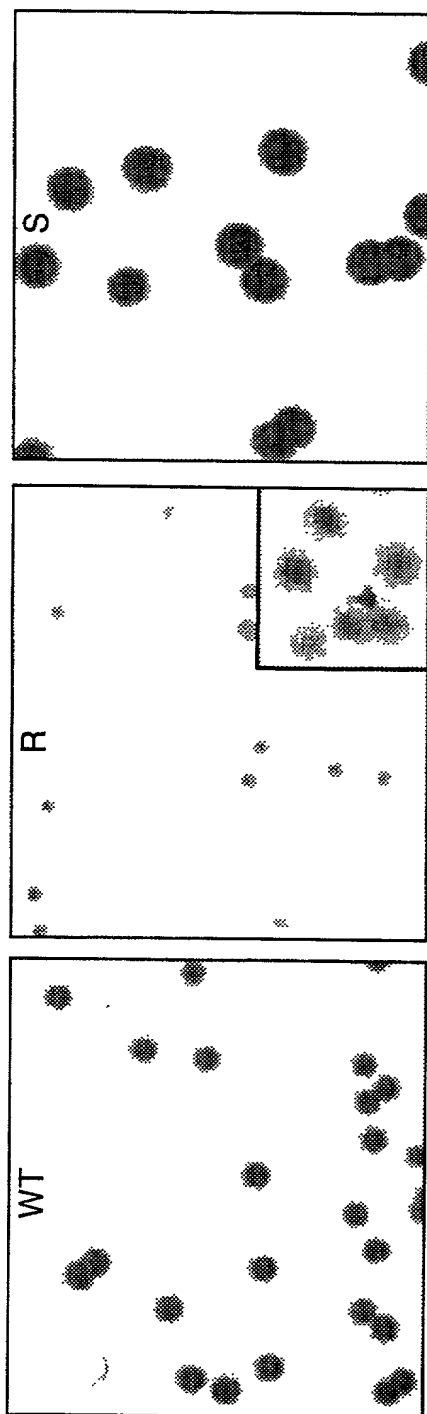


Fig.1b.

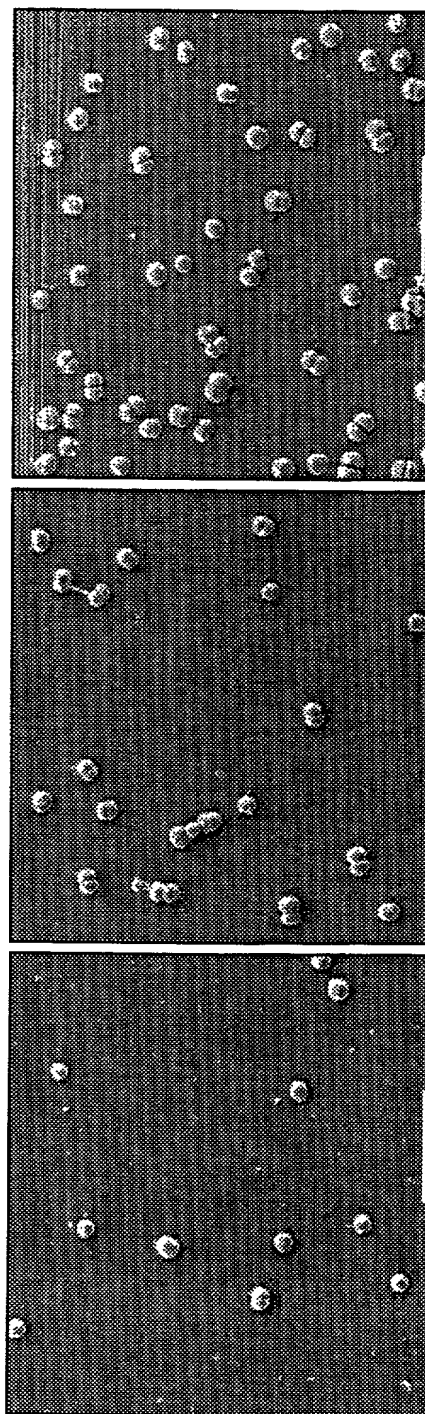


Fig.1c.

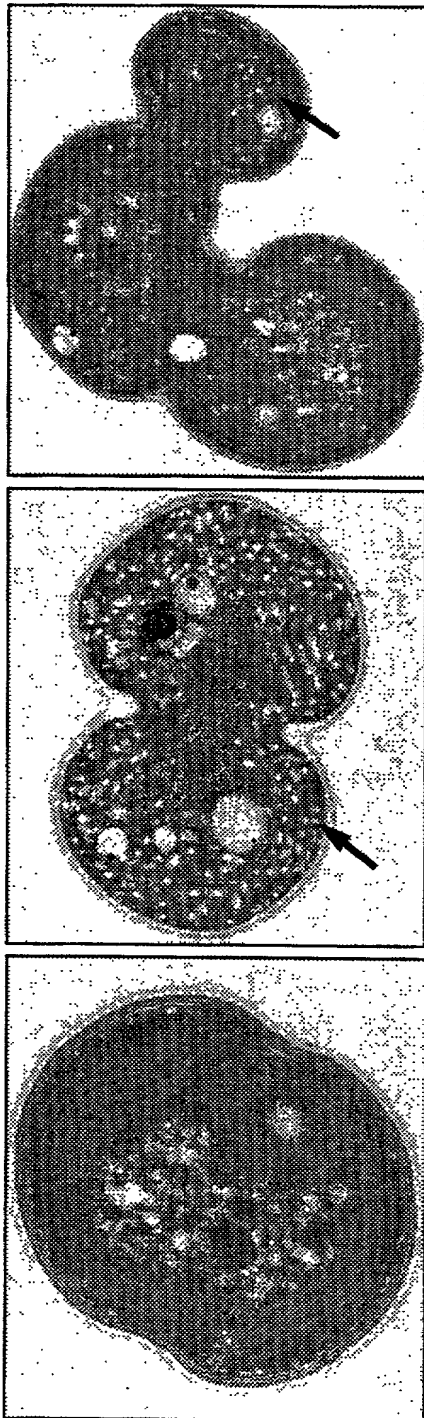
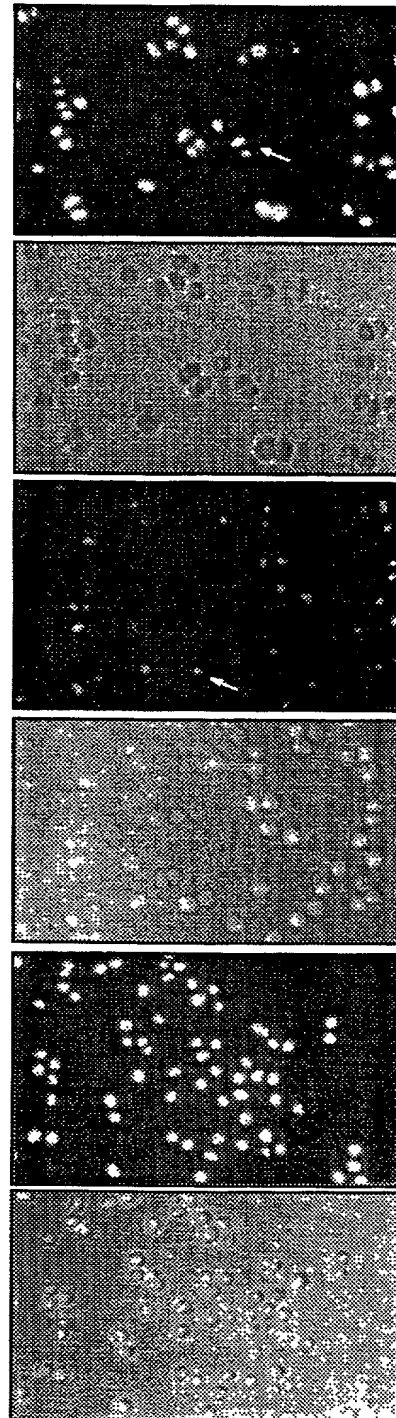


Fig.1d.



3/4

Fig.2.

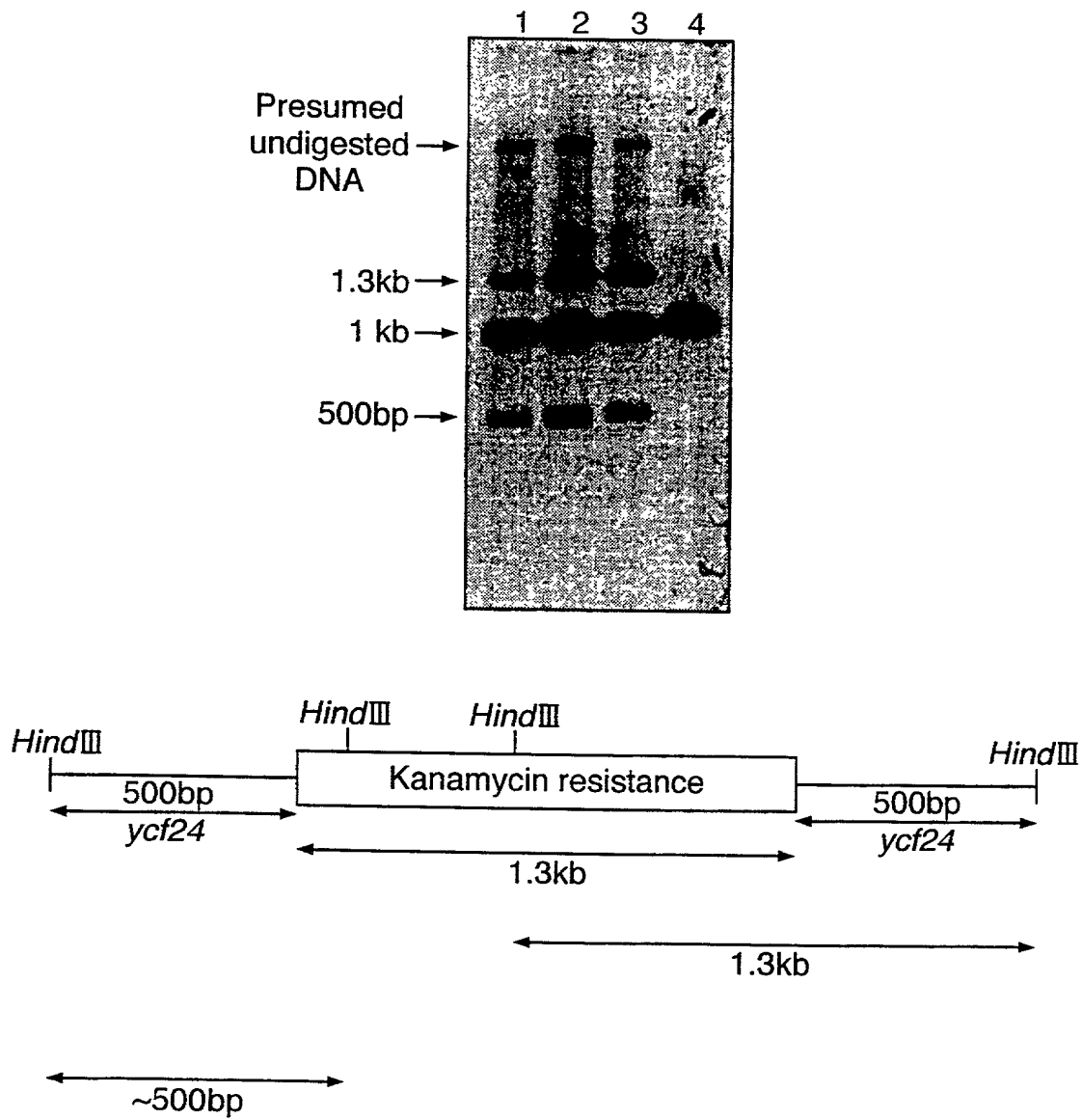


Fig.3a.

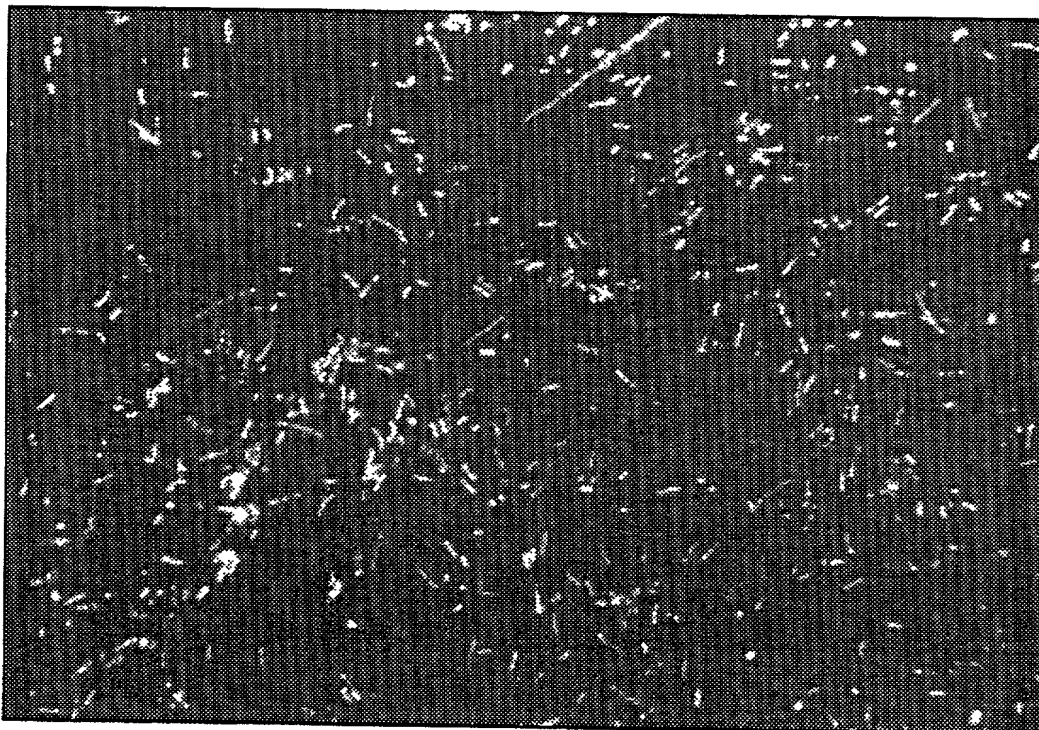
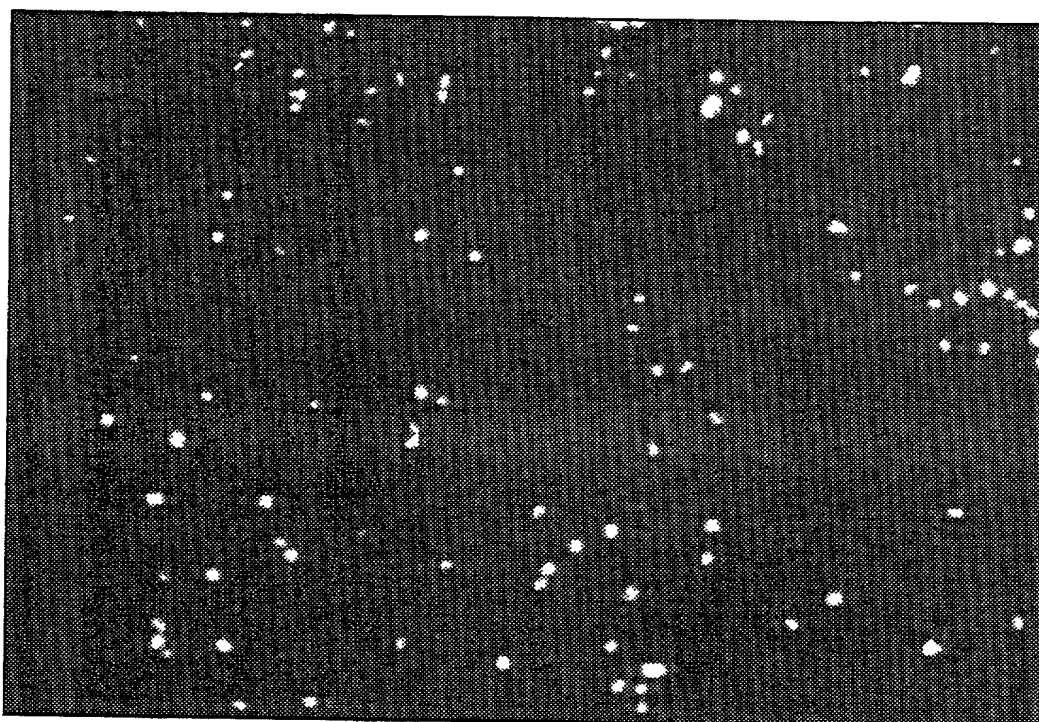


Fig.3b.



RULE 63 (37 C.F.R. 1.63)
**DECLARATION AND POWER OF ATTORNEY
 FOR PATENT APPLICATION
 IN THE UNITED STATES PATENT AND TRADEMARK OFFICE**

As a below named inventor, I hereby declare that my residence, post office address and citizenship are as stated below next to my name, and I believe I am the original, first and sole inventor (if only one name is listed below) or an original, first and joint inventor (if plural names are listed below) of the subject matter which is claimed and for which a patent is sought on the invention entitled INHIBITORS OF YCF24 AS ANTIMALARIALS

the specification of which (check applicable box(es)).

☐ is attached hereto

☐ was filed on

☒ was filed as PCT international application No. PCT/GB 99/03180 on 22nd September 1999

and (if applicable to U.S. or PCT application) was amended on

I hereby state that I have reviewed and understand the contents of the above identified specification, including the claims, as amended by any amendment referred to above. I acknowledge the duty to disclose information which is material to the examination of this application in accordance with 37 C.F.R. 1.56(a). I hereby claim foreign priority benefits under 35 U.S.C. 119/365 of any foreign application(s) for patent or inventor's certificate listed below and have also identified below any foreign application for patent or inventor's certificate having a filing date before that of the application on which priority is claimed or, if no priority is claimed, before the filing date of this application.

Prior Foreign Application(s):

Application Number	Country	Day/Month/Year Filed
9820658.4	United Kingdom	22 nd September 1998

I hereby claim the benefit under 35 U.S.C. 120/365 of all prior United States and PCT international applications listed above or below and, insofar as the subject matter of each of the claims of this application is not disclosed in such prior applications in the manner provided by the first paragraph of 35 U.S.C. 112, I acknowledge the duty to disclose material information as defined in 37 C.F.R. 1.56(a) which occurred between the filing date of the prior applications and the national or PCT international filing date of this application.

Prior U.S./PCT Application(s):

Application Serial No.	Day/Month/Year Filed	Status patented, pending, abandoned
PCT/GB 99/03180	22 nd September 1999	

I hereby declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true, and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

And I hereby appoint Nixon & Vanderhye P.C., 1100 North Glebe Road, 8th Floor, Arlington, Virginia 22201-4714, telephone number (703) 816-400 (to whom all communications are to be directed), and the following attorneys thereof (of the same address) individually and collectively my attorneys to prosecute this application and to transact all business in the Patent and Trademark Office connected therewith and with the resulting patent: Arthur R. Crawford, 25327; Larry S. Nixon, 25640; Robert A. Vanderhye, 27076; James T. Hosmer, 30184; Robert W. Farris, 31352; Richard G. Besha, 22770; Mark E. Nusbaum, 32348; Michael J. Keenan, 32106; Bryan H. Davidson, 30251; Stanley C. Spooner, 27393; Leonard C. Mitchard, 29009; Duane M. Byers, 33363; Paul J. Henon, 33626; Jeffry H. Nelson, 30481; John R. Lastova, 33149; H. Warren Burnam, Jr., 29366; Thomas E. Byrne, 32205.

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First Middle Initial Family Name Citizenship

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2) Inventor's Signature Conrad Mullineaux Date 8/5/01

Inventor's Name (typed) Conrad William MULLINEAUX British
First Middle Initial Family Name Citizenship

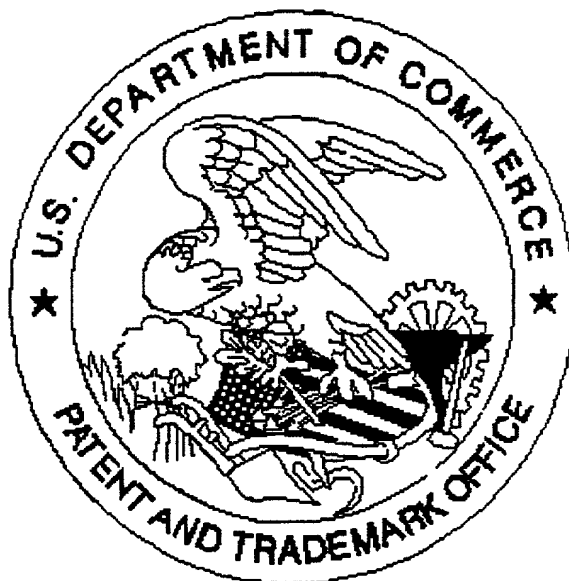
Residence (City) Barnet (State/Foreign Country) Hertfordshire United Kingdom GBX

Post Office Address 112 Mays Lane, Barnet, Hertfordshire EN5 2LS United Kingdom Zip Code EN5 2LS

FOR ADDITIONAL INVENTORS, check box [X] and attach sheet with same information and signature and date for each.

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